An Investigation of Compression Techniques to Speed up Mutation Testing

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Mutation Testing

An actively investigated field since 1970s

• Main idea: small syntactic changes \rightarrow test suite quality

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Our paper: speed up

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Challenged by Gopinath et al. (ICSE 2016,TR 2017):

No practical advantage over pure random sampling



647 KB



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19 KB



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- Necessity
- Sufficiency

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Program: 1 int fun(int a, int b){ 2 int c; 3 if(a>0) 4 c=a+b; 5 else 6 c=a-b; 7 return abs(c); 8 }

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- Necessity \rightarrow weak mutation
- Sufficiency



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Overall methodology



Mutant clustering

Based on weak mutation

• Overlapped grouping

elements are only grouped together if they are identical

• FCA-based grouping

Formal Concept Analysis; convey binary relations

	t ₁	t_2	t3	t ₄
m_1	0	1	0	0
m ₂	1	0	0	0
m ₃	0	0	0	1
m4	0	0	1	0
m_5	1	1	0	0
m ₆	1	1	0	0

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• Cluster-based selection (CS)

randomly chooses one mutant from each cluster

• CS + mutation operator type (mop)

divide each cluster into partitions by mutation operator types and then random selection

• CS + mutation location (mloc)

	mop	mloc
m_1	1	Line5
m ₂	1	Line5
m3	2	Line5
m ₄	2	Line6

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■ 20 open-source projects (Java): 397K+ LOC,

- 2K+ tests, 166K+ mutants
- 6+2 methods:
 - overlap
- pure random • fca

- overlap+mop fca+mop weak mutation
- overlap+mloc
 fca+mloc

Research questions:

- **RQ1:** How accurate are different compression techniques?
- **RQ2:** How do compression techniques perform in terms of speed-up?
- **RQ3:** What is the trade-off between accuracy and speed-up?

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baselines

RQ1: accuracy performance

• Absolute error

$$AE(C, T) = | strong_M(C, T) - estimated_M(C, T) |$$
 (1)

• Results:



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Absolute error

$$AE(C, T) = | strong_M(C, T) - estimated_M(C, T) |$$
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method comparison

RQ2: Speed-up performance

Speed-up



Results:



(2)

RQ2: Speed-up performance

Speed-up



(2)

RQ3: Trade-off

Speed-up v.s. absolute error (mean)



RQ3: Trade-off

Speed-up v.s. prediction accuracy (mean)



RQ3: Trade-off

Speed-up v.s. prediction accuracy (mean)



Sum-up



• Mutation compression v.s. random sampling:



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- Mutation compression: speed-up \uparrow and accuracy \uparrow
- Killable mutant results outperform overall mutation score



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- Mutation compression v.s. random sampling:
 - $\bullet\,$ Mutation compression: speed-up \uparrow and accuracy $\uparrow\,$
 - Killable mutant results outperform overall mutation score
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- Future work:
 - Combining mutation operator and mutation location
 - Exploring other compression techniques
 - Applying to test-data generation